PU48040-C SEQ LIST

SEQUENCE LISTING

```
<110> SmithKline Beecham Corporation
<120> ErbB4 Co-Crystal
<130> PU4804WO
<150> 60/441,443
<151> 2003-01-21
<160> 2
<170> FastSEQ for Windows Version 4.0
<210> 1
<211> 364
<212> PRT
<213> homo sapien
<400> 1
Met Lys Lys Gly His His His His His Gly Leu Glu Thr Glu Leu
1 5 10 15
Val Glu Pro Leu Thr Pro Ser Gly Thr Ala Pro Asn Gln Ala Gln Leu 20 25 30
Arg Ile Leu Lys Glu Thr Glu Leu Lys Arg Val Lys Val Leu Gly Ser
                             40
Gly Ala Phe Gly Thr Val Tyr Lys Gly Ile Trp Val Pro Glu Gly Glu
50 55 60
Thr Val Lys Ile Pro Val Ala Ile Lys Ile Leu Asn Glu Thr Thr Gly
                     70
Pro Lys Ala Asn Val Glu Phe Met Asp Glu Ala Leu Ile Met Ala Ser
                                       90
Met Asp His Pro His Leu Val Arg Leu Leu Gly Val Cys Leu Ser Pro
                                  105
                                                         110
Thr Ile Gln Leu Val Thr Gln Leu Met Pro His Gly Cys Leu Leu Glu
115 120 125
                              120
Tyr Val His Glu His Lys Asp Asn Ile Gly Ser Gln Leu Leu Leu Asn
                          135
    130
                                                140
Trp Cys Val Gln Ile Ala Lys Gly Met Met Tyr Leu Glu Glu Arg Arg
145 150 155 160
Leu Val His Arg Asp Leu Ala Ala Arg Asn Val Leu Val Lys Ser Pro
165 170 175
Asn His Val Lys Ile Thr Asp Phe Gly Leu Ala Arg Leu Leu Glu Gly
180 185 190
Asp Glu Lys Glu Tyr Asn Ala Asp Gly Gly Lys Met Pro Ile Lys Trp
_ 195 _ _ 200 205
Met Ala Leu Glu Cys Ile His Tyr Arg Lys Phe Thr His Gln Ser Asp
                          215
                                                220
Val Trp Ser Tyr Gly Val Thr Ile Trp Glu Leu Met Thr Phe Gly Gly
                     230
                                           235
Lys Pro Tyr Asp Gly Ile Pro Thr Arg Glu Ile Pro Asp Leu Leu Glu
                 245
                                       250
Lys Gly Glu Arg Leu Pro Gln Pro Pro Ile Cys Thr Ile Asp Val Tyr 260 265 _ 270
Met Val Met Val Lys Cys Trp Met Ile Asp Ala Asp Ser Arg Pro Lys
        275
                              280
Phe Lys Glu Leu Ala Ala Glu Phe Ser Arg Met Ala Arg Asp Pro Gln 290 295 300
Arg Tyr Leu Val Ile Gln Gly Asp Asp Arg Met Lys Leu Pro Ser Pro
                    310
                                           315
                                           Page 1
```

PU48040-C SEQ LIST

Asn Asp Ser Lys Phe Phe Gln Asn Leu Leu Asp Glu Glu Asp Leu Glu
325

Asp Met Met Asp Ala Glu Glu Tyr Leu Val Pro Gln Ala Phe Asn Ile
340

Pro Pro Pro Ile Tyr Thr Ser Arg Ala Arg Ile Asp
355

360

<210> 2 <211> 966 <212> DNA <213> homo sapien

atgaaaaaag gtcatcatca tcatcatcat ggtttggaaa cagagttggt ggaaccatta 60 actcccagtg gcacagcacc caatcaagct caacttcgta ttttgaaaga aactgagctg 120 aagaggggtaa aagtccttgg ctcaggtgct tttggaacgg tttataaagg tatttgggta 180 cctgaaggag aaactgtgaa gattcctgtg gctattaaga ttcttaatga gacaactggt 240 cccaaggcaa atgtggagtt catggatgaa gctctgatca tggcaagtat ggatcatcca 300 cacctagtcc ggttgctggg tgtgtgtctg agcccaacca tccagctggt tactcaactt 360 atgccccatg gctgctgtt ggagtatgtc cacgagcaca aggataacat tggatcacaa 420 ctgctgctta actggtgtg ccagatagct aagggaatga tgtacctgga agaaagacga 480 ctcgttcatc gggatttggc agcccgtaat gtcttagtga aatctccaaa ccatgtgaaa 540 ggaggaaaga tgccaattaa atggatggct ctggagtgt tacatacagg aactgttggag ctatggagt actatatgg aactgatgac catcgagga acgtttggag ctatggagt actatatgg aactgatgac ctttggagga 720 aaaccctatg atggaattcc aacgcgagaa atccctgatt tattagagaa aggagaacgt 780 atggagcccc aaagatacc aacgtagacc taaatttaag gaactggctg ctgagtttc aaggatggct 900 cgagaccctc aaagatacc agttatcag ggtgatgatc gtatgaagct tcccagtcca 960 aattga